



FIG. 1

>SEQID#1_SGP037
ATGTCAACAGCTGCCTTAATTACTTTGGTCAGAAGTGGTGGGAACCAAGGTGAGAAGGAGAGTGTCTAAGCTCCGGCTGCTGCAGGACGACA
GGCGGGTGACACCCACAGTCCACTTCAGAGCCTAGGTGTTCTCGGTTTGACCCAGATGGTAGTGGGAGTCCAGCTACCTGGGACAA
TTTTGGGATCTGGGATAACCGCATTGATGAGCCAATTCTGCTGCCACCCAGCAATTAAGTATGGCAAGCCAATCCCCAAAATCAGCTTGGAAAATG
TGGGGTGGCCTCACAGATTGGCAACGGAAGAGAAATGAAGATCGGTTTGACTTCGCTCAGCTGACAGATGAGGTCTCTGACTTTGCAGTGTA
TGATGGACACGGTGGACCTGCAGCAGCTGATTTCTGTCTATACCCACATGGAGAAATGTATTATGGATTTGCTTCTTAAGGAGAAGAACTTGGAA
ACTCTGTTGACCTTGGCTTTTCTAGAAATAGATAAAGCCTTTTCGAGTCAATGCCCGCTGCTGCTGATGCAACTCTTCTGACCTCTGGGACTACT
GCAACAGTAGCCCTATTGCGAGATGGTATTGAACCTGGTTGTAGCCAGTGTGGGGACAGCCGGGCTATTTGTGTAGAAAAGGAAAACCCATGA
AGCTGACCATTGACCATACTCCAGAAAGAAAAGATGAAAAGAAAGGATCAAGAAATGTGGTGGTTTTGTAGCTTGGAAATAGTTGGGGCAGC
CTCAGGTAATGGCAGGCTTGCAATGACAAGAAGTATTGGAGATTTGGACCTTAAGACCAGTGGTGTCTATAGCAGAACCTGAACCTAAGAGGAT
TAAGTTACATCATGCTGATGACAGCTTCTGGTCTCCACACAGATGGAATTAACCTCATGGTGAATAGTCAAGAGATTTGTGACTTTGTCAATC
AGTGCCATGATCCCAACGAAGCAGCCCATGCGGTGACTGAACAGGCAATACAGTACGGTACTGAGGATACAGTACTGCAGTAGTAGTGCCTTT
TGGTGCCTGGGGAAAAATAAGAAGCTCTGAATCAACTTCTCATTACAGAGAAGCTTTGCCTCCAGTGGACGATGGGCCTGA

FIG. 2

>SEQID#2_SGP037
MSTAALITLVRSGNQVRRRVLLSSRLLQDDRRVTPTCHSSTSEPRCSRFPDGGSPATWDFGTDNRIDEPIILLPPSIKYGKPIPKISLENVGCASQIG
KRKENEDRFQALTDDEVLYFAVYDGHGGPAAADFCHTHMEKCIDMLLPKEKNLETLTLAFLFIDKAFSSHARLSADATLLTSGTTATVALLRDGIEL
VVASVGDSRAILCRKGKPNKLTIDHTPERKDEKERIKKCGGFVAVNSLGQPHVNGRLAMTRSIGDLDLKTSGVIAEPETKRKIKLHHADDSFLVLTDDGI
NFMVNSQEI CDFVNQCHDPNEAAHAVTEQAIOYGTEDNSTAVVVPFGAWGKYKNSINFSSRSFSSGRWA